

<u>→</u>

## 1 MWILALSLEGSFANVFSEDLHSS

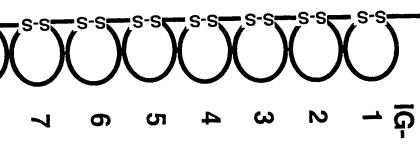
Signal Peptide

- 24 LYFVNASLQEVVFASTTGTLVPCPAAGIPPVTLRWYLATGEEIYDVPGIRHVHPNGTLQIFPFPPSSF STLIHDNTYYCTAENPSGKIRSQDVHIKAVLREPY
- 127  ${\tt TVRVEDQKTMRGNVAVFKCIIPSSVEAYITVVSWEKDTVSLVSGSRFLITSTGALYIKDVQNEDGLYN}$ YR**C**ITRHRYTGETRQSNSARLFVSDPANSAP
- 226 SILDGEDHRKAMAGQRVELPCKALGHPEPDYRWLKDNMPLELSGREQKTVTGLLIENIRPSDSGSYVC EVSNRYGTAKVIGRLYVKQPLKA
- 317 TISPRKVKSSVGSQVSLSCSVTGTEDQELSWYRNGEILNPGKNVRITGINHENLIMDHMVKSDGGAYQ CFVRKDKLSAQDYVQVVLEDGTPKI
- 410 ISAFSEKVVSPAEPVSLMCNVKGTPLPTITWTLDDDPILKGGSHRISQMITSEGNVVSYLNISSSQVR DGGVYR**C**TANNSAGVVLYQARINVRGPAS
- 507  ${\tt IRPMKNITAIAGRDTYIH}{\tt CRVIGYPYYSIKWYKNSNLLPFNHRQVAFENNGTLKLSDVQKEVDEGEYT}$ CNVLVQPQLSTSQSVHVTVKVPPFIQPFE
- 604 FPRESIGQRVFIPCVVVSGDLPITITWQKDGRPIPGSLGVTIDNIDFTSSLRISNLSLMHNGNYTCIA RNEAAAVEHQSQLIVRVPPKFVVQPR
- 698 DQDGIYGKAVILNCSAEGYPVPTIVWKFSKGAGVPQFQPIALNGRIQVLSNGSLLIKHVVEEDSGYYL CKVSNDVGADVSKSMYLTVKIPAMITS

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793 YPNTTLATQGQKKEMSCTAHGEKPIIVRWEKEDRIINPEMARYLVSTKEVGEEVISTLQILPTVREDS GFFSCHAINSYGEDRGIIQLTVQEPPD

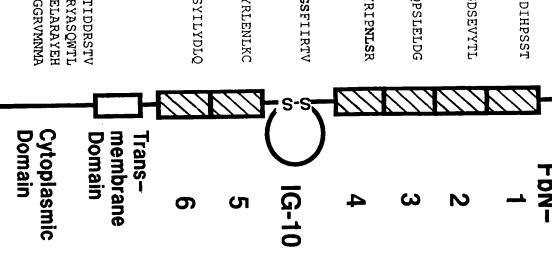
## FIG. 2A





- 888 PPEIEIKDVKARTITLRWTMGFDGNSPITGYDIECKNKSDSWDSAQRTKDVSPQLNSATIIDIHPSST YSIRMYAKNRIGKSEPSNELTITADEAA
- 984 PDGPPQEVHLEPISSQSIRVTWKAPKKHLQNGIIRGYQIGYREYSTGGNFQFNIISVDTSGDSEVYTL DNLNKFTQYGLVVQACNRAGTGPSSQEIITTTLED
- 1087 VPSYPPENVQAIATSPESISISWSTLSKEALNGILQGFRVIYWANLMDGELGEIKNITTTQPSLELDG LEKYTNYSIQVLAFTRAGDGVRSEQIFTRTK
- 1186 EDVPGPPAGVKAAAASASMVFVSWLPPLKLNGIIRKYTVFCSHPYPTVISEFEASPDSFSYRIP**NLS**R NRQYSVWVVAVTSAGRGNSSEIITVEPL
- 1282 AKAPARILTESGTVTTPWMKDIVLPCKAVGDPSPAVKWMKDSNGTPSLVTIDGRRSIFSNGSFIIRTV KAEDSGYYSCIANNNWGSDEIILNLQ
- 1376 VQVPPDQPRLTVSKTTSSSITLSWLPGDNGGSSIRGYILQYSEDNSEQWGSFPISPSERSYRLENLKC GTWYKFTLTAQNGVGPGRISEIIEAKTL
- 1472 GKEPQFSKEQELFASINTTRVRLNLIGWNDGGCPITSFTLEYRPFGTTVWTTAQRTSLSKSYILYDLQ EATWYELQMRVCNSAGCAEKQANFATLNYDGSTIPPLIKSVVQNEEGLTTNEGLK
- 1595 MLVTISCILVGVLLLFVLLLVV
- 1617 RRRREQRLKRLRDAKSLAEMLMSKNTRTSDTLSKQQQTLRMHIDIPRAQLLIEERDTMETIDDRSTV AKMEEQLRHAKFTITECFISDTSSEQLTAGTNEYTDSLTSSTPSESGICRFTASPPKPQDGGRVMNMA NRPHPTISAHTLTTDWRLPTPRAAGSVDKESDSYSVSPSQDTDRARSSMVSTESASSTYEELARAYEH VPKAIGQVTSYICLHTLEWTFC LLTDADFGEAAKQKSLTVTHTVHYQSVSQATGPLVDVSDARPGTNPTTRRNAKAGPTARNRYASQWTL

FIG. 2B





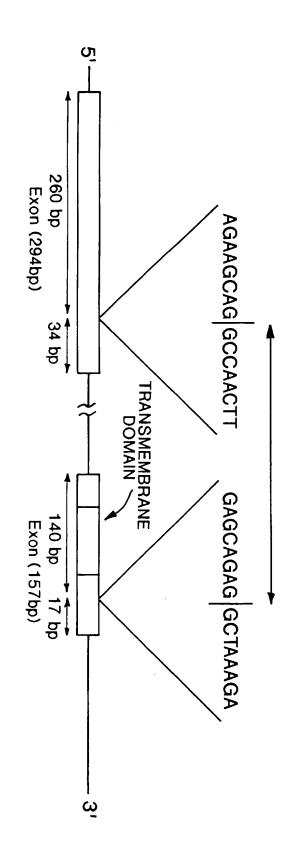
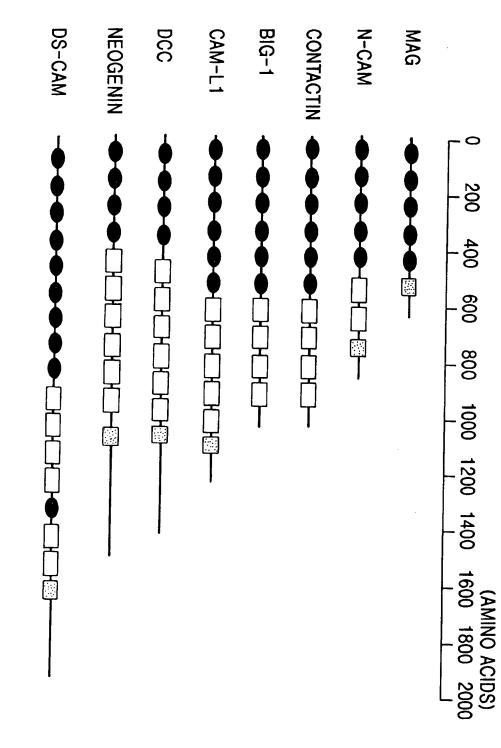


FIG. 3





Ig TYPE-C2 DOMAIN

FIBRONECTIN TYPE-III DOMAIN

TRANSMEMBRANE DOMAIN

FIG. 4